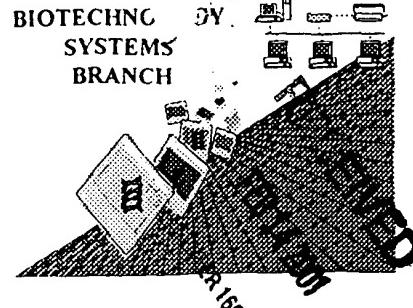


D Srivastava



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/424,487A

Source: 1653

Date Processed by STIC: 1-17-01

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
 - 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**
- FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.**

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1 825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO).

Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

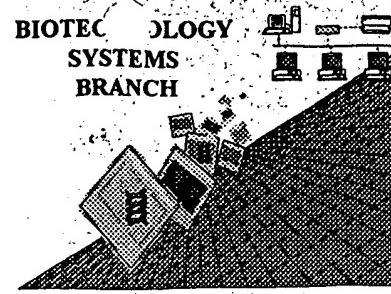
Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

Hoppe
Robinson

new

RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/424,487A

Source: 1653

Date Processed by STIC: 1/17/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:
<http://www.uspto.gov/web/offices/pac/checker>

1653

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/424,487A

DATE: 01/17/2001
TIME: 12:20:34

Input Set : A:\CHOO1.APP.txt
Output Set: N:\CRF3\01172001\I424487A.raw

3 <110> APPLICANT: CHOO, Yen
4 KLUG, Aaron
5 ISALAN, Mark
7 <120> TITLE OF INVENTION: Nucleic Acid Binding Proteins
9 <J30> FILE REFERENCE: 71278/264975
11 <140> CURRENT APPLICATION NUMBER: US 09/424,487A
C--> 12 <141> CURRENT FILING DATE: 2000-02-29
14 <150> PRIOR APPLICATION NUMBER: GB 9710809.6
15 <151> PRIOR FILING DATE: 1997-05-23
17 <150> PRIOR APPLICATION NUMBER: PCT/GB98/01512
18 <151> PRIOR FILING DATE: 1998-05-26
20 <160> NUMBER OF SEQ ID NOS: 17
22 <170> SOFTWARE: PatentIn Ver. 2.1
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 264
26 <212> TYPE: DNA
27 <213> ORGANISM: Artificial Sequence
29 <220> FEATURE:
30 <221> NAME/KEY: CDS
31 <222> LOCATION: (1)..(264)
33 <220> FEATURE:
34 <223> OTHER INFORMATION: Description of Artificial Sequence: encoding
35 nucleic acid binding proteins
37 <400> SEQUENCE: 1
38 gca gaa gag aag cct ttt cag tgt cga atc tgc atg cgt aac ttc agc 48
39 Ala Glu Glu Lys Pro Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser
40 1 5 10 15
42 gat cgt act act ctt acc cgc cac acg agg acc cac aca ggc gag aag 96
43 Asp Arg Thr Thr Leu Thr Arg His Thr Arg Thr His Thr Gly Glu Lys
44 20 25 30
46 cct ttt cag tgt cga atc tgc atg cgt aac ttc agc agg agc gat aac 144
47 Pro Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser Arg Ser Asp Asn
48 35 40 45
50 ctt acg aga cac cta agg acc cac aca ggc gag aag cct ttt cag tgt 192
51 Leu Thr Arg His Leu Arg Thr His Thr Gly Glu Lys Pro Phe Gln Cys
52 50 55 60
54 cga atc tgc atg cgt aac ttc agg caa gct gat cat ctt caa gag cac 240
55 Arg Ile Cys Met Arg Asn Phe Arg Gln Ala Asp His Leu Gln Glu His
56 65 70 75 80
58 cta aag acc cac aca ggc gag aag 264
59 Leu Lys Thr His Thr Gly Glu Lys
60 85
63 <210> SEQ ID NO: 2
64 <211> LENGTH: 88
65 <212> TYPE: PRT
66 <213> ORGANISM: Artificial Sequence
68 <220> FEATURE:

Does Not Comply
Corrected Diskette Needed

pp. 2, 3, 4

A RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/424,487A

DATE: 01/17/2001
TIME: 12:20:34

Input Set : A:\CHOO1.APP.txt
Output Set: N:\CRF3\01172001\I424487A.raw

69 <223> OTHER INFORMATION: Description of Artificial Sequence: encoding
70 nucleic acid binding proteins
72 <400> SEQUENCE: 2
73 Ala Glu Glu Lys Pro Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser
74 1 5 10 15
76 Asp Arg Thr Thr Leu Thr Arg His Thr Arg Thr His Thr Gly Glu Lys
77 20 25 30
79 Pro Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser Arg Ser Asp Asn
80 35 40 45
82 Leu Thr Arg His Leu Arg Thr His Thr Gly Glu Lys Pro Phe Gln Cys
83 50 55 60
85 Arg Ile Cys Met Arg Asn Phe Arg Gln Ala Asp His Leu Gln Glu His
86 65 70 75 80
88 Leu Lys Thr His Thr Gly Glu Lys
89 85
93 <210> SEQ ID NO: 3
94 <211> LENGTH: 18
95 <212> TYPE: PRT
96 <213> ORGANISM: Artificial Sequence
98 <220> FEATURE:
99 <223> OTHER INFORMATION: Description of Artificial Sequence: Nucleic acid
100 binding protein
102 <220> FEATURE:
103 <221> NAME/KEY: BINDING
104 <222> LOCATION: (1)..(18) /
105 <223> OTHER INFORMATION: where X is any amino acid
107 <400> SEQUENCE: 3
W--> 108 Xaa Cys Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa His Xaa Xaa
109 1 5 10 15
W--> 111 Xaa His
115 <210> SEQ ID NO: 4
116 <211> LENGTH: 21
117 <212> TYPE: PRT
118 <213> ORGANISM: Artificial Sequence
120 <220> FEATURE:
121 <223> OTHER INFORMATION: Description of Artificial Sequence: where X is any
122 amino acid
124 <220> FEATURE:
125 <221> NAME/KEY: BINDING
126 <222> LOCATION: (1)..(21)
128 <400> SEQUENCE: 4 <223>
W--> 129 Xaa Cys Xaa Xaa Cys Xaa Xaa Phe Xaa Xaa Xaa Xaa Xaa Leu Xaa Xaa
130 1 5 10 15
W--> 132 His Xaa Xaa Xaa His
133 20
136 <210> SEQ ID NO: 5
137 <211> LENGTH: 26
138 <212> TYPE: PRT
139 <213> ORGANISM: Artificial Sequence

where is source of
the artificial sequence?

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/424,487A

DATE: 01/17/2001
TIME: 12:20:34

Input Set : A:\CHOOL.APP.txt
Output Set: N:\CRF3\01172001\I424487A.raw

```

207      probability of any of the four bases at DNA position 3
209 <220> FEATURE:
210 <221> NAME/KEY: BINDING
211 <222> LOCATION: (1)..(9)
213 <400> SEQUENCE: 8
W--> 214 Gly Asn Xaa Xaa Cys Gly Gly Cys Gly
215     1           5
218 <210> SEQ ID NO: 9
219 <211> LENGTH: 9
220 <212> TYPE: PRT
221 <213> ORGANISM: Artificial Sequence
223 <220> FEATURE:
224 <223> OTHER INFORMATION: Description of Artificial Sequence: where X
225 denotes a known combination of the two bases at
226 DNA positions 4X and 5X
228 <220> FEATURE:
229 <221> NAME/KEY: BINDING
230 <222> LOCATION: (1)..(9)
232 <400> SEQUENCE: 9
W--> 233 Gly Cys Xaa Xaa Cys Gly Gly Cys Gly
234     1           5
237 <210> SEQ ID NO: 10
238 <211> LENGTH: 28
239 <212> TYPE: PRT
240 <213> ORGANISM: Artificial Sequence
242 <220> FEATURE:
243 <223> OTHER INFORMATION: Description of Artificial Sequence: Zinc finger
244 binding protein
246 <220> FEATURE:
247 <221> NAME/KEY: BINDING
248 <222> LOCATION: (1)..(28)
250 <400> SEQUENCE: 10
251 Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser Asp Arg Ser Ser Leu
252     1           5           10          15
254 Thr Arg His Thr Arg Thr His Thr Gly Glu Lys Pro
255     20          25
258 <210> SEQ ID NO: 11
259 <211> LENGTH: 28
260 <212> TYPE: PRT
261 <213> ORGANISM: Artificial Sequence
263 <220> FEATURE:
264 <223> OTHER INFORMATION: Description of Artificial Sequence: Zinc finger
265 binding protein
267 <220> FEATURE:
268 <221> NAME/KEY: BINDING
269 <222> LOCATION: (1)..(28)
271 <400> SEQUENCE: 11
272 Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser Asp Arg Ser His Leu
273     1           5           10          15

```

"Xaa" cannot denote
a base.
Description of artificial
Sequence?

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/424,487A

DATE: 01/17/2001
TIME: 12:20:34

Input Set : A:\CH001.APP.txt
Output Set: N:\CRF3\01172001\I424487A.raw

275 Thr Arg His Thr Arg Thr His Thr Gly Glu Lys Pro
276 20 25
279 <210> SEQ ID NO: 12
280 <211> LENGTH: 27
281 <212> TYPE: PRT
282 <213> ORGANISM: Artificial Sequence
284 <220> FEATURE:
285 <223> OTHER INFORMATION: Description of Artificial Sequence: Zinc finger
286 binding protein
288 <220> FEATURE:
289 <221> NAME/KEY: BINDING
290 <222> LOCATION: (1)..(27)
292 <400> SEQUENCE: 12
293 Phe Gln Cys Arg Ile Cys Met Arg Asn Phe Ser Asp Arg Ser Asn Leu
294 1 5 10 15
296 Thr Arg His Thr Arg Thr His Thr Gly Glu Lys
297 20 25
300 <210> SEQ ID NO: 13
301 <211> LENGTH: 9
302 <212> TYPE: PRT
303 <213> ORGANISM: Artificial Sequence
305 <220> FEATURE:
306 <223> OTHER INFORMATION: Description of Artificial Sequence: Zinc finger
307 binding protein
309 <220> FEATURE:
310 <221> NAME/KEY: BINDING
311 <222> LOCATION: (1)..(9)
313 <400> SEQUENCE: 13
314 Ala Gly Ala Gly Ala Gly Cys Thr Cys
315 1 5
318 <210> SEQ ID NO: 14
319 <211> LENGTH: 8
320 <212> TYPE: PRT
321 <213> ORGANISM: Artificial Sequence
323 <220> FEATURE:
324 <223> OTHER INFORMATION: Description of Artificial Sequence: 8bp
325 palindromic sequence which is bound and cleaved by
326 the restriction endonuclease NotI
328 <220> FEATURE:
329 <221> NAME/KEY: BINDING
330 <222> LOCATION: (1)..(8)
332 <400> SEQUENCE: 14
333 Gly Cys Gly Gly Cys Cys Gly Cys
334 1 5
337 <210> SEQ ID NO: 15
338 <211> LENGTH: 9
339 <212> TYPE: PRT
340 <213> ORGANISM: Artificial Sequence
342 <220> FEATURE:

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/424,487A

DATE: 01/17/2001
TIME: 12:20:35

Input Set : A:\CHOOL.APP.txt
Output Set: N:\CRF3\01172001\I424487A.raw

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:108 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:111 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:129 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:132 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:194 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:214 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:233 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9



Creation date: 08-25-2003

Indexing Officer: TLAM2 - THY LAM

Team: OIPEBackFileIndexing

Dossier: 09424487

Legal Date: 03-28-2001

No.	Doccode	Number of pages
1	CRFL	7

Total number of pages: 7

Remarks:

Order of re-scan issued on